

Comparison of MetAP2 Homologues (mouse = SEQ ID NO:13; rat = SEQ ID NO:17;  
human = SEQ ID NO:12; yeast = SEQ ID NO:14)

mouse	1	15	16	30	31	45	46	60	61	75	76	90
MAGVEQAASFGGHLN	GDLDPPDREEGTST	AEEAARRRRRRRRR	KGAVSAVQQLDKES	GALVDEVAKQLESA	LEEKERDDDDDDG	90						
rat	MAGVEAASFGGHLN	RDLDPPDREEGTST	AEEAARRRRRRRRR	KGAVSAVQQLDKES	GTSVDEVAKQLERA	LEEKERDDDDDDG	90					
human	MAGVEEVAASGSHLN	GDLDPPDREEGAAT	AEEAARRRRRRRRR	KGPSAAGEQEPDKES	GASVDEVAKQLERA	LEEKERDDDDDDG	90					
yeast	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	-----	38
mouse	91	105	106	120	121	135	136	150	151	165	166	180
DADGATGKKKKKKKK	KRGPVQTDPSVPI	CDLYPVGVPKQEC	EYPTQDGRTAAMRT	TSEKKALDQASEEI	WDFREAAEAHRQVR	180						
rat	DGDGAAGKKKKKKKK	KRGPVQTDPSVPI	CDLYPVGVPKQEC	EYPTQDGRTAAMRT	TSEKKALDQASEEI	WDFREAAEAHRQVR	180					
human	DGDGATGKKKKKKKK	KRGPVQTDPSVPI	CDLYPVGVPKQEC	EYPTQDGRTAAMRT	TSEKKALDQASEEI	WDFREAAEAHRQVR	180					
yeast	ESKKKKKKKKKKKK	N-----	VKKI	ELLPDGKYPGAMM	DYHDFNLQRTTDEE	SRYLKRDLERA--EH	WNVKRGAEIHRVR	116				
mouse	181	195	196	210	211	225	226	240	241	255	256	270
KYVMSWIKPGMTMIE	ICEKLEDCSRKLKE	NGLNAG-----	LA	FPTGCSLNCCAHT	PNAGDTTVLQYDDIC	KIDFGTHISGRIDC	263					
rat	KYVMSWIKPGMTMIE	ICEKLEDCSRKLKE	NGLNAG-----	LA	FPTGCSLNCCAHT	PNAGDTTVLQYDDIC	KIDFGTHISGRIDC	263				
human	KYVMSWIKPGMTMIE	ICEKLEDCSRKLKE	NGLNAG-----	LA	FPTGCSLNCCAHT	PNAGDTTVLQYDDIC	KIDFGTHISGRIDC	263				
yeast	RAIKDRIVPGMKLMD	IADMIENTTRKYTGA	ENLLAMEDPKSQSIG	FPTGLSLNHCAHFT	PNAGDKTVLKYEDEM	KVDYGVQVNGNIIDS	206					
mouse	271	285	286	300	301	315	316	330	331	345	346	360
AFTVTENPKYDILLT	AVKDATNTGICAGI	DVRLCDVGEALQEV	ESYEVEIDGKTYQVK	PIRLNNGHSIGPYRI	HAGKTVPIVKGGEAT	353						
rat	AFTVTENPKYDILLT	AVKDATNTGICAGI	DVRLCDVGEALQEV	ESYEVEIDGKTYQVK	PIRLNNGHSIGPYRI	HAGKTVPIVKGGEAT	353					
human	AFTVTENPKYDILLT	AVKDATNTGICAGI	DVRLCDVGEALQEV	ESYEVEIDGKTYQVK	PIRLNNGHSIGPYRI	HAGKTVPIVKGGEAT	353					
yeast	AFTVSFDQYDNLILA	AVKDATYTGICKEAGI	DVRLTDIGEALQEV	ESYEVEINGEYTYQVK	PCRNLCGHSIAPYRI	HGKSVPIVKNGDTT	296					
mouse	361	375	376	390	391	405	406	420	421	435	436	450
RMEEGEVAIETFGS	TGKGVHDMDCSHY	MKNEDVGHPVIRLPR	TKHLNVINENEGTL	AFCRRLDRLGESKY	LMALKNLCDLGI	443						
rat	RMEEGEVAIETFGS	TGKGVHDMDCSHY	MKNEDVGHPVIRLPR	TKHLNVINENEGTL	AFCRRLDRLGESKY	LMALKNLCDLGI	443					
human	RMEEGEVAIETFGS	TGKGVHDMDCSHY	MKNEDVGHPVIRLPR	TKHLNVINENEGTL	AFCRRLDRLGESKY	LMALKNLCDLGI	443					
yeast	KMEEGEHAETFGS	TGRGYVTAGGEVSHY	ARSAEDHQVMPITLDS	AKNLKTIIDRNFGTL	PFCRRYLDRLGQEKY	LFALNNLVHRHGLVQD	386					
mouse	451	465	466	480								
YPPLCDIKGSYTAQF	EHTILLRPTCKEVVS	RGDDY--	478									
rat	YPPLCDIKGSYTAQF	EHTILLRPTCKEVVS	RGDDY--	478								
human	YPPLCDIKGSYTAQF	EHTILLRPTCKEVVS	RGDDY--	478								
yeast	YPPLNDIPGSYTAQF	EHTILLHAHKEVVS	KGDDY--	421								

Figure 1

Title: Dominant Negative Variants fo Mehtionine  
Aminopeptidase  
Inventor(s): Chang et al.  
Appln. No. 09/943,123  
Docket # 66153/45004

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## MetAP2

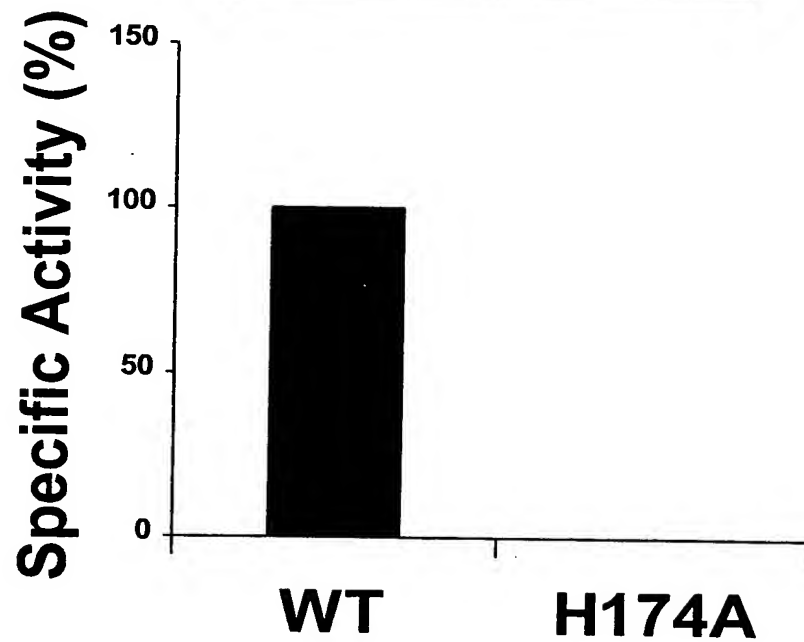
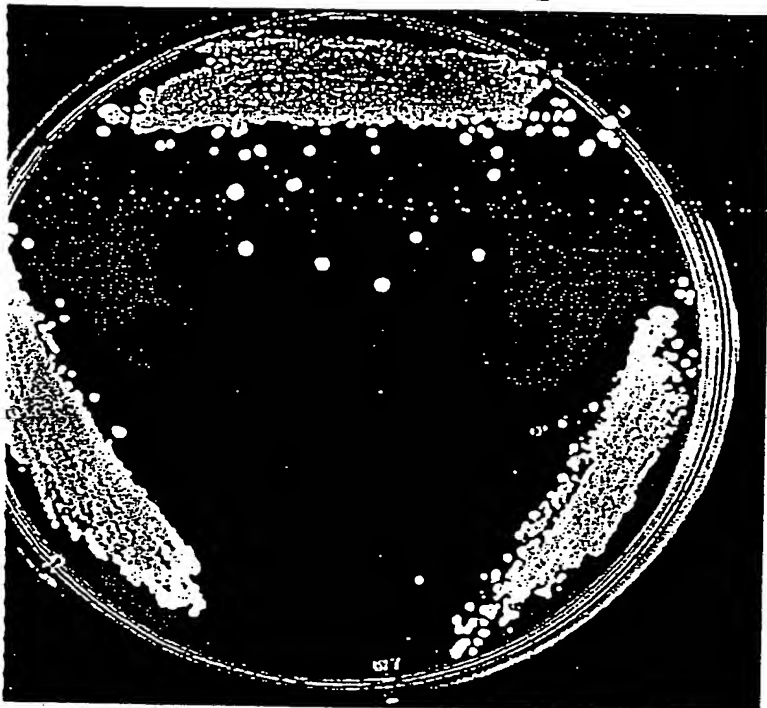
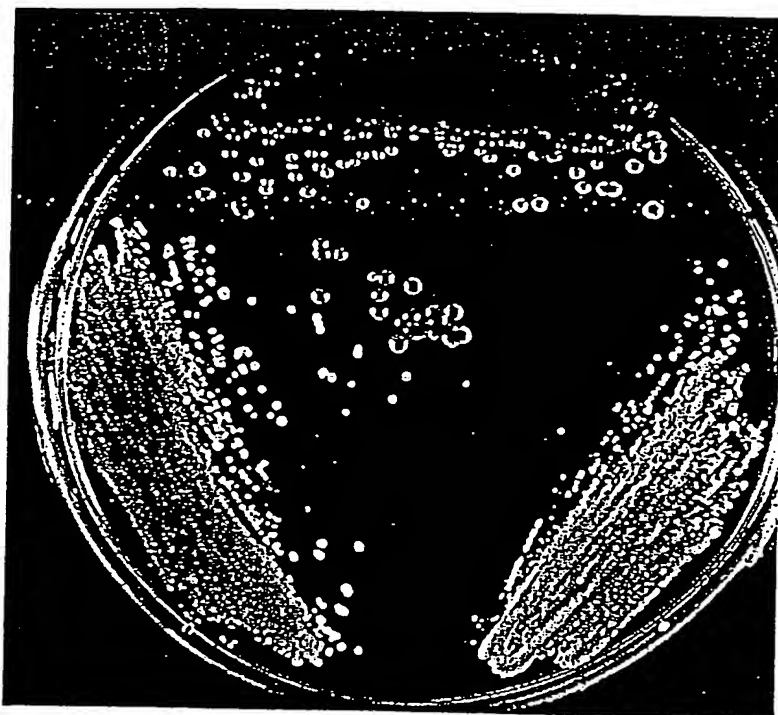


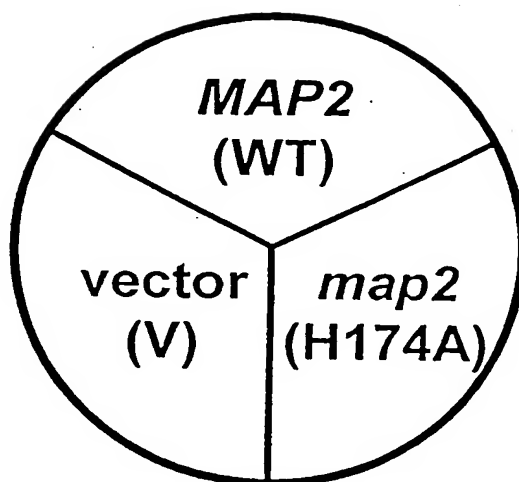
Figure 2



**A. Glucose**



**B. Galactose**



**FIGURE 3**

Title:	Dominant Negative Variants fo Mehtionine Aminopeptidase
Inventor(s):	Chang et al.
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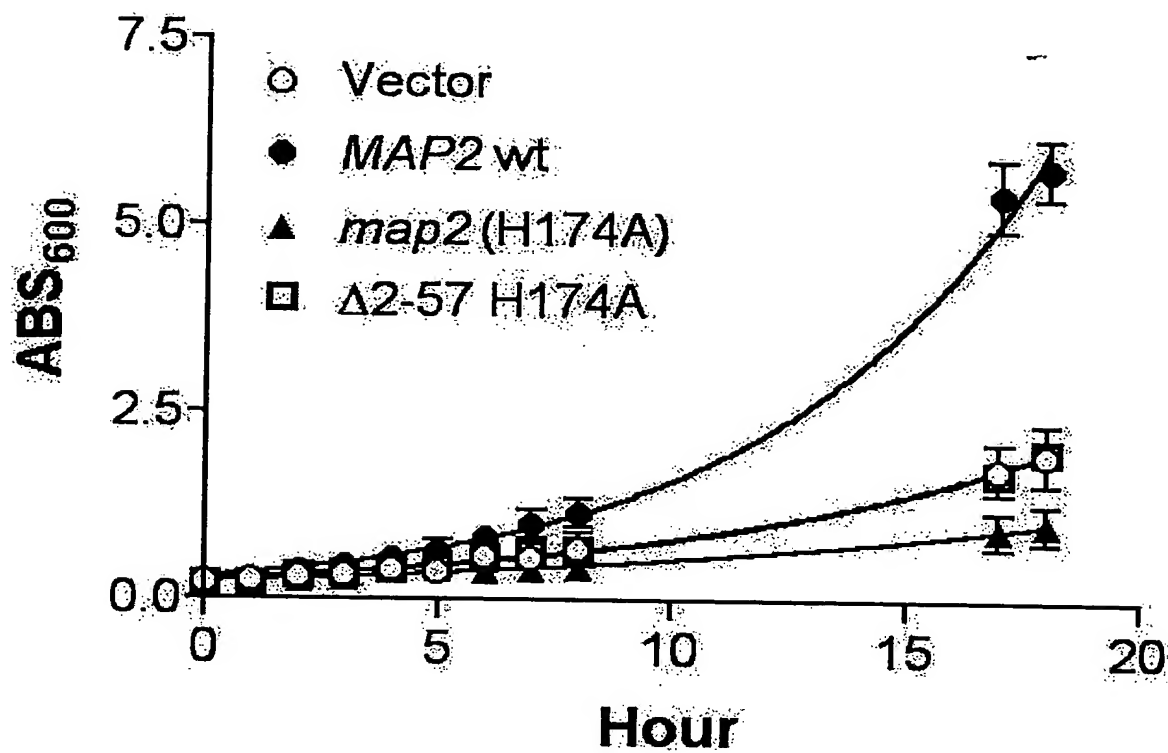
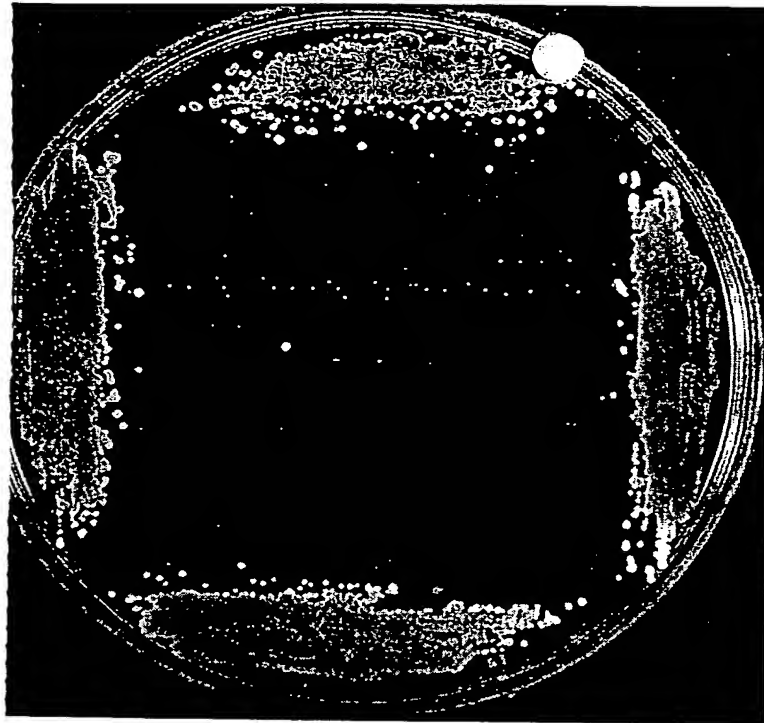
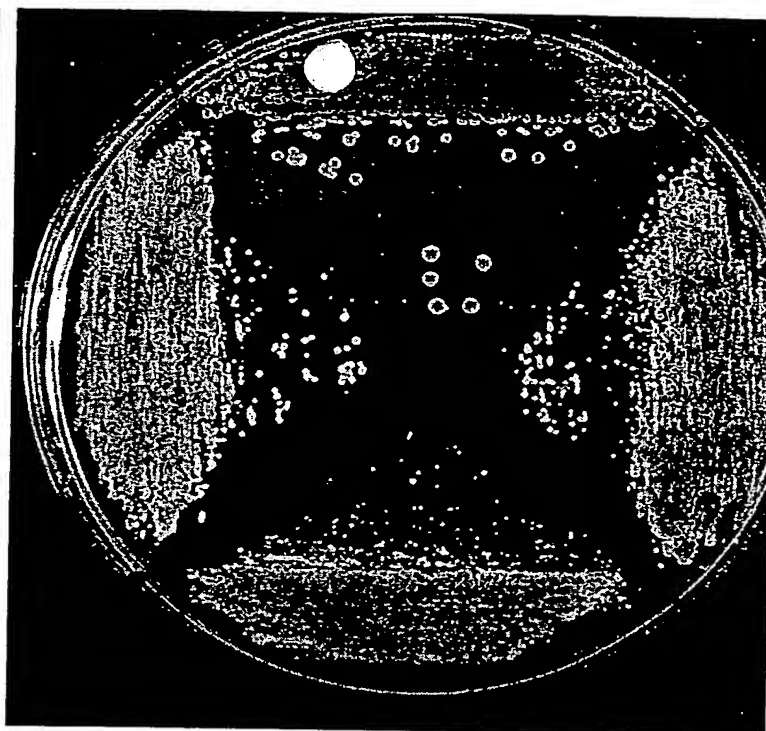


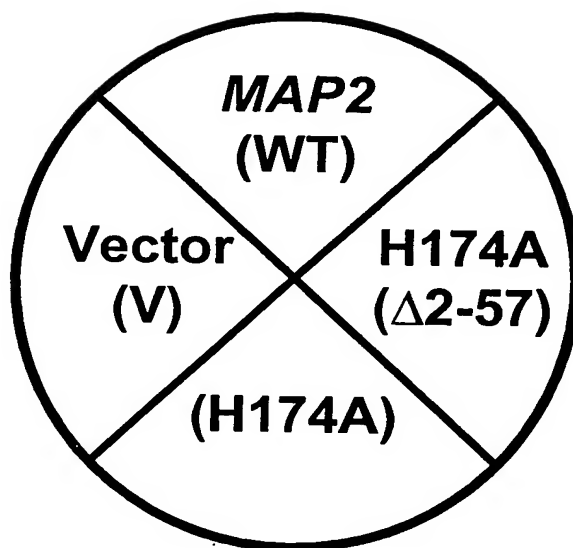
Figure 4



**A. Glucose**



**B. Galactose**



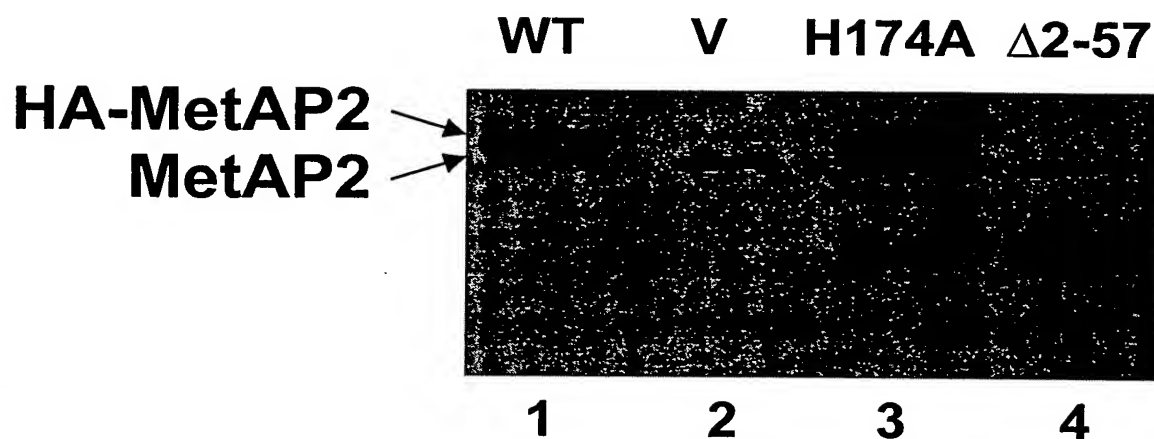
H174A-MetAP2 requires N-terminal residues 2-57 for inhibition of *map1Δ* growth under the GAL1 promoter.

**Figure 5**

Title:	Dominant Negative Variants fo Mehtionine Aminopeptidase
Inventor(s):	Chang et al.
Appln. No.	09/943,123
Docket #	66153/45004

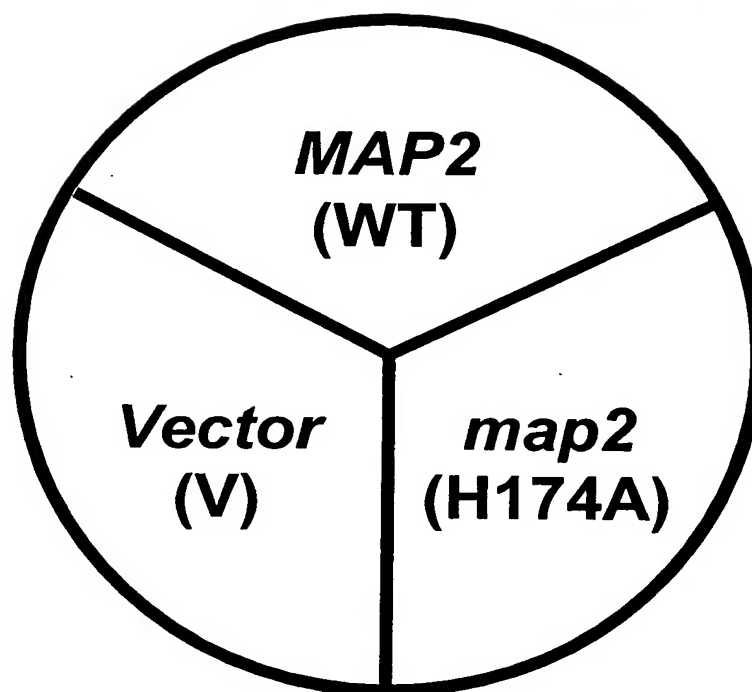
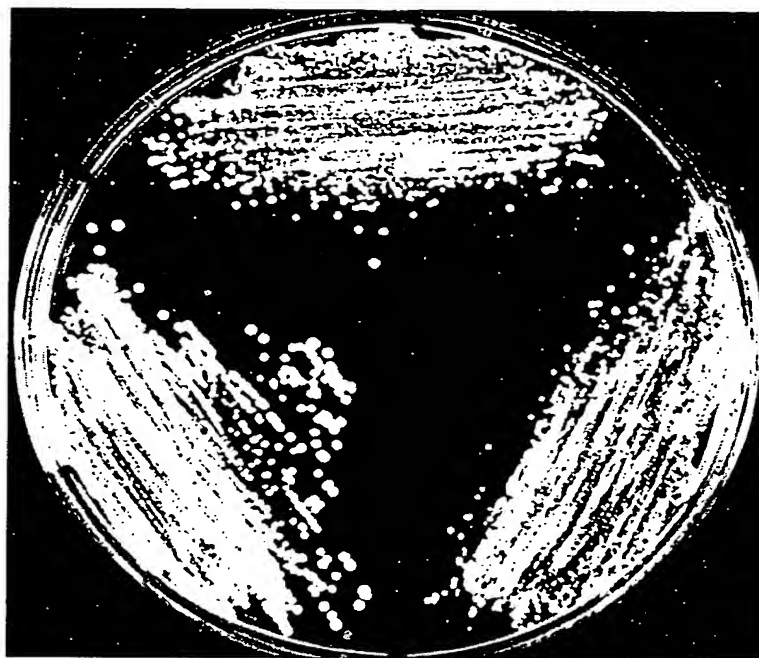
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Inventor(s): Chang et al.  
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The steady state levels of each MetAP2 construct are comparable. Immunoblot comparison of HA-MetAP2 wt, HA-MetAP2 H174A, and MetAP2 Δ2-57 H174A steady state levels in map1Δ.

Figure 6



Overexpression of H174A-MetAP2 under the GPD promoter does not inhibit the growth of map2Δ

Figure 7

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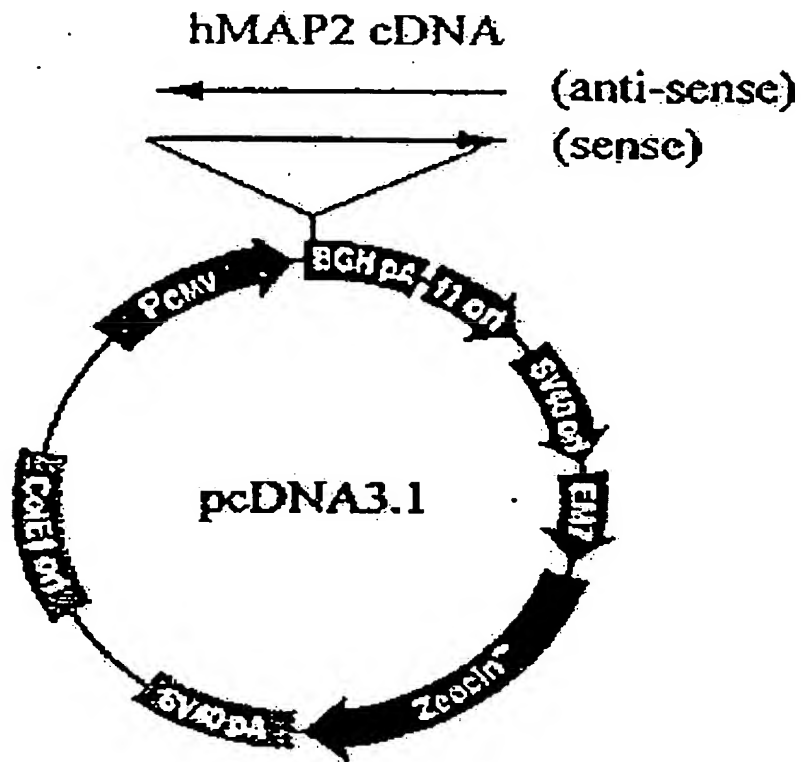


Figure 8



Title: Dominant Negative Variants fo Mehtionine  
 Amino peptidase  
 Inventor(s): Chang et al.  
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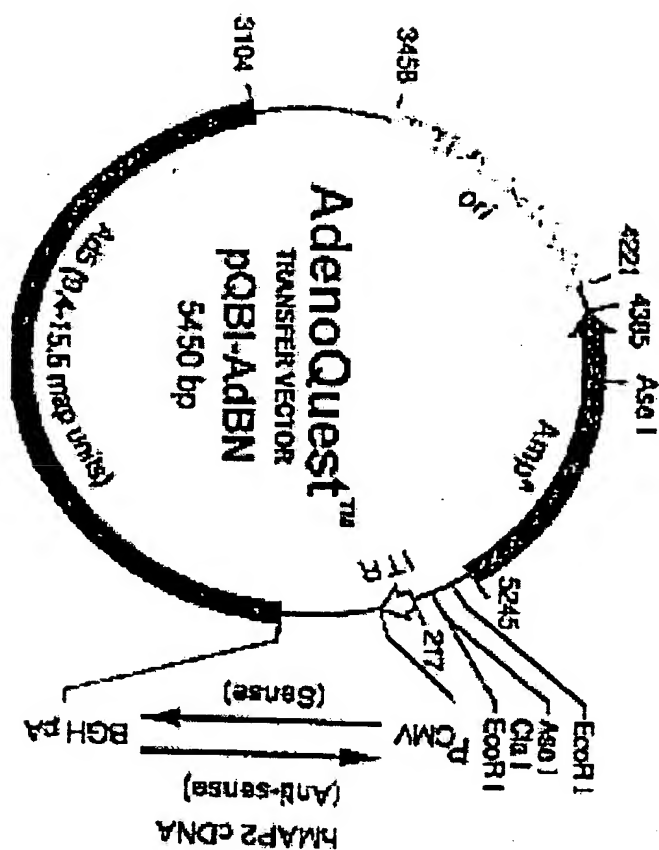


Figure 9

Title: Dominant Negative Variants fo Mehtionine  
Aminopeptidase  
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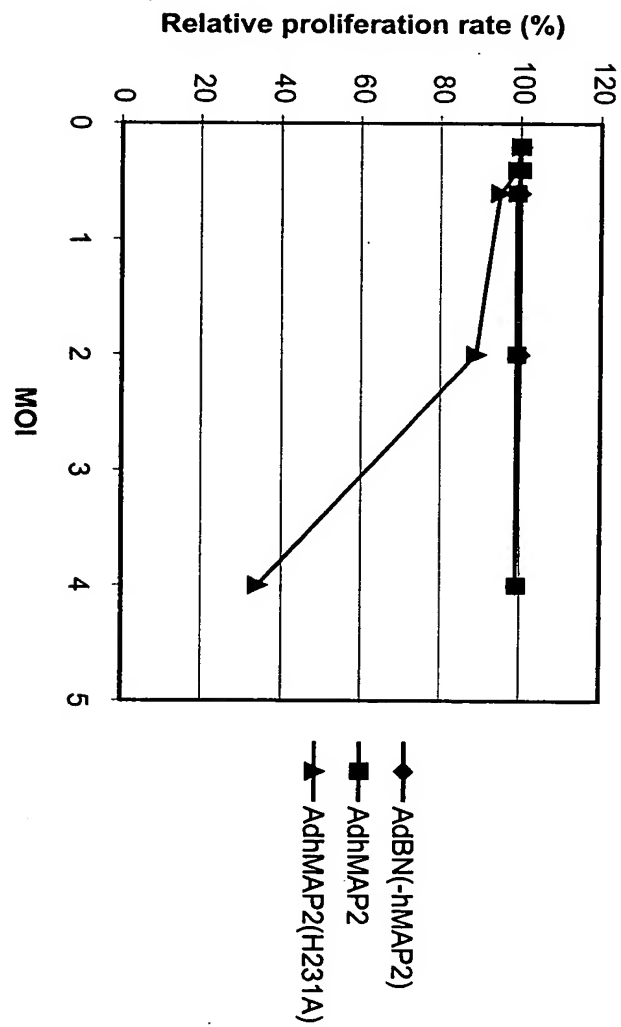


Figure 10

Title: Dominant Negative Variants fo Mehtionine  
 Aminoamidase  
 Inventor(s): Chang et al.  
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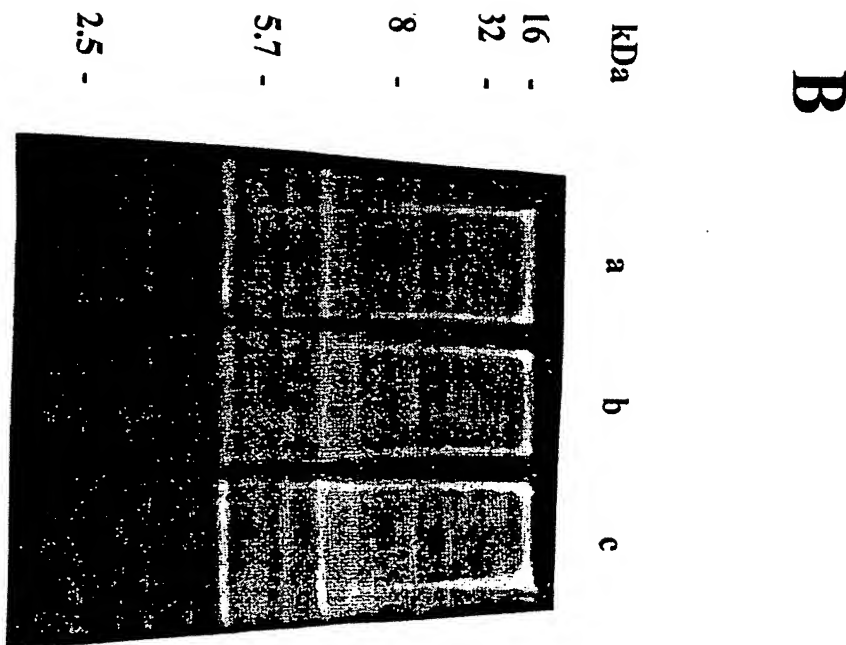
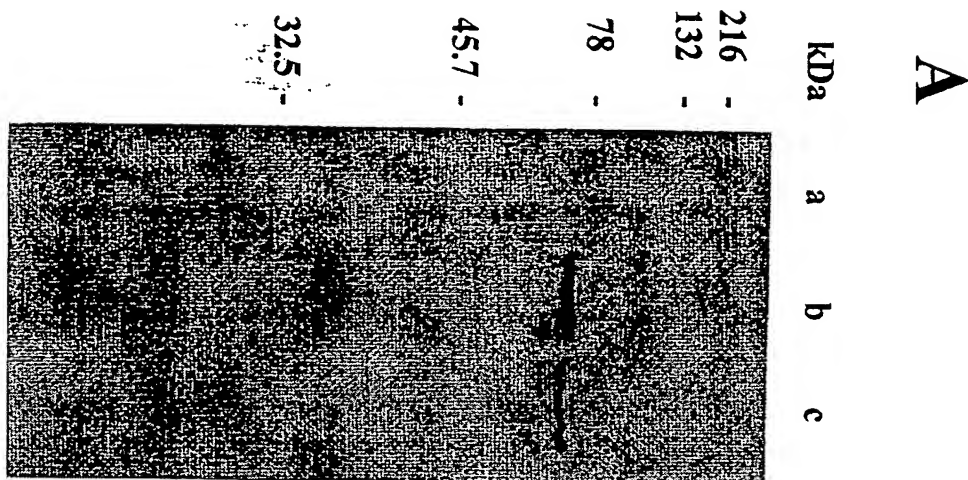


Figure 11